

Please amend the title by replacing the current title with the following:

3-PHOSPHOINOSITIDE-DEPENDENT PROTEIN KINASE

Please amend the paragraph on page 39, line 18 to page 40, line 1 as follows:

Figure 17. Evidence the PDK1 and DSTPK61 possess a Pleckstrin-Homology domain. Sequence alignment of PH domains that have a known tertiary structure with the putative PH domains of PDK1 and DSTPK61 (SEQ ID NOS:30-35). Alignment was done using the program AMPS (Barton & Sternberg (1990) *J. Mol. Biol.* 212, 389-402) and formatted using AMAS (Livingstone & Barton (1993) *Comp. Appl. Bio. Sci.* 9, 745-756) with some manual adjustment to ensure coincidence of secondary structure regions. Regions containing conserved residues are coloured green and the invariant tryptophan (Trp-535) is coloured red. General regions of secondary structure are denoted by the blue (beta strand) and green (alpha helix) bands. The Pleckstrin-Homology sequences are those found in human Pleckstrin, PLS; human spectrin, SPC; human dynamin DYN, and rat phospholipase C- δ , PLC δ . Numbering is based on PDK1 sequence (SEQ ID NO:1) (Fig. 10).